

HYS-38CIP sequence listing
SEQUENCE LISTING

<110> Deder, Douglas
Yamazaki, Victoria
Asundi, Vinod
Liu, Chenghua
Tang, Y. Tom
Drmanac, Radoje T.

<120> Methods of Therapy and Diagnosis Using Insulin-like Growth Factor Binding Protein-like Polypeptides and Polynucleotides

<130> HYS-38CIP

<140> Not Yet Assigned
<141> 2002-02-27

<150> 09/784,748
<151> 2001-02-14

<150> 09/649,167
<151> 2000-08-23

<150> 09/540,217
<151> 2000-03-31

<160> 14

<170> PatentIn version 3.1

<210> 1
<211> 375

HYS-38CIP sequence listing

<212> DNA

<213> Homo sapiens

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tcaccggggc gcaggtgggc ctgtcctgtg aagtgagggc tgtgcctacc ccagtcatca 180
cgtggagaaa ggtcacgaag tccctgagg gcacccaagc actggaggag ctgcctgggg 240
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<212> DNA

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atgttctaag tcattttcac tattttcacac ccattttacg agatatttga ggtggcttat 180
aagacctgtta gaaaaaagaa gaaaaatacg taaatggagg aaaccaggaa aagagcaaaa 240
gaagagtagg gacatactta gatgagcagt agaatccctg gtatattctg cacacatctc 300
cctctgagct tcttagcatg caaagacaag agctgtgaac atgaagggtgt gtccatgaga 360
tgaaaagacc agttgtgttt tgggctgga gggatattt cctctgtatt ctttttagaaa 420
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<211> 375

<212> DNA

<213> Homo sapiens

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HYS-38CIP sequence listing

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cgtggagaaa ggtcacgaag tccctgagg gcacccaagc actggaggag ctgcctgggg	240
accatgtcaa tatagctgtc caagtgcgag gggcccttc tgaccatgag gccacggcct	300
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<212> DNA

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ccgggatctc ggcgcgtcgcgagatgcgct gctgcgttgc ctgcctggaa gccgaggcg	300
cgagctgcgg gggccgcgc ggcggcgct gtggcccg cctggatgc gcgagccagg	360
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gcggctccga cggcgctcg taccggcg tctgcgtgcgc gtcggcaca	480
cgccccggcgc gcaccccggt cacctgcaca aggcgcgcga cggccgtgc gagttcggttc	540
ctatcaactcg ttttataac tgcttcctc agccgttaat tcacaggcaa ttctttgt	600
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aacacaccat taactttaaa gaaatctcg agggatttgg gaagatttt tcattccagc	780
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tgctggatgc cagggtggct gagttgtgt ccaatgcgc tcctgtggc gtcgttgc	900
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ccacctactc ctactccagc accctttctc cttcacaggt gtttctccta atacatctct	1200

HYS-38CIP sequence listing

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<212> DNA

<213> Homo sapiens

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Met Pro Arg Leu Ser Leu Leu Leu Pro Leu Leu
1 5 10

ctt ctg ctg ctg ctg ccg ctg ctg ccg ccg ctg tcc ccg agc ctc ggg
 Leu Leu Leu Leu Leu Pro Leu Leu Pro Pro Leu Ser Pro Ser Leu Gly
 15 20 25 159

atc cgc gac gtg ggc ggc cgg cgc ccc aag tgt ggt ccg tgc cgg cca 207
 Ile Arg Asp Val Gly Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro
 30 35 40

gag ggc tgc ccg gcg cct gcg ccc tgc ccg gcg ccc ggg atc tcg gcg 255
 Glu Gly Cys Pro Ala Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala
 45 50 55

60 65 70 75 303
 ctc gag gag tgc ggc tgc tgc gcc cgc tgc ctg gga gcc gag ggc gcg
 Leu Asp Glu Cys Cys Gly Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala
 60 65 70 75 303

agc tgc ggg ggc cgc gcc ggc ggg cgc tgt ggc ccc ggc ctg gta tgc 351
 Ser Cys Gly Gly Arg Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys
 80 85 90

gcg agc cag gcc gct ggg gca gcg ccc gag ggc acc ggg ctc tgc gtg 399
 Ala Ser Gln Ala Ala Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val
 95 100 105

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 Cys Ala Gln Arg Gly Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro
 110 115 120

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Ser Val Cys Ala Leu Arg Leu Arg Ala Arg His Thr Pro Arg Ala His
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HYS-38CIP sequence listing

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160 165 170	
gtg ggc ctg tcc tgt gaa gtg agg gct gtg cct acc cca gtc atc acg	639
Val Gly Leu Ser Cys Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr	
175 180 185	
tgg aga aag gtc acg aag tcc cct gag ggc acc caa gca ctg gag gag	687
Trp Arg Lys Val Thr Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu	
190 195 200	
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Leu Pro Gly Asp His Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro	
205 210 215	
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Ser Asp His Glu Ala Thr Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys	
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gag gat gag ggt gtg tac cag tgc cat gca gcc aac atg gtg gga gag	831
Glu Asp Glu Gly Val Tyr Gln Cys His Ala Ala Asn Met Val Gly Glu	
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Ala Glu Ser His Ser Thr Val Thr Val Leu Asp Leu Ser Lys Tyr Arg	
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<212> PRT
<213> Homo sapiens

<400> 6
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Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro Glu Gly Cys Pro Ala
35 40 45

HYS-38CIP sequence listing

Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala Leu Asp Glu Cys Gly
50 55 60

Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala Ser Cys Gly Gly Arg
65 70 75 80

Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys Ala Ser Gln Ala Ala
85 90 95

Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val Cys Ala Gln Arg Gly
100 105 110

Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro Ser Val Cys Ala Leu
115 120 125

Arg Leu Arg Ala Arg His Thr Pro Arg Ala His Pro Gly His Leu His
130 135 140

Lys Ala Arg Asp Gly Pro Cys Glu Phe Ala Pro Val Val Val Val Pro
145 150 155 160

Pro Arg Ser Val His Asn Val Thr Gly Ala Gln Val Gly Leu Ser Cys
165 170 175

Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr Trp Arg Lys Val Thr
180 185 190

Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu Leu Pro Gly Asp His
195 200 205

Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro Ser Asp His Glu Ala
210 215 220

Thr Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys Glu Asp Glu Gly Val
225 230 235 240

Tyr Gln Cys His Ala Ala Asn Met Val Gly Glu Ala Glu Ser His Ser
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Thr Val Thr Val Leu Asp Leu Ser Lys Tyr Arg Ser Phe His Phe Pro
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Ala Pro Asp Asp Arg Met
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HYS-38CIP sequence listing

<211> 837

<212> DNA

<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

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<211> 27

<212> PRT

<213> Homo sapiens

HYS-38CIP sequence listing

<400> 9

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<210> 10

<211> 251

<212> PRT

<213> Homo sapiens

<400> 10

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35 40 45

Ser Cys Gly Gly Arg Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys
50 55 60

Ala Ser Gln Ala Ala Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val
65 70 75 80

Cys Ala Gln Arg Gly Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro
85 90 95

Ser Val Cys Ala Leu Arg Leu Arg Ala Arg His Thr Pro Arg Ala His
100 105 110

Pro Gly His Leu His Lys Ala Arg Asp Gly Pro Cys Glu Phe Ala Pro
115 120 125

Val Val Val Val Pro Pro Arg Ser Val His Asn Val Thr Gly Ala Gln
130 135 140

Val Gly Leu Ser Cys Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr
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HYS-38CIP sequence listing

Trp Arg Lys Val Thr Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu
165 170 175

Leu Pro Gly Asp His Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro
180 185 190

Ser Asp His Glu Ala Thr Ala Trp Ile Leu Ile Asn Pro Leu Arg Lys
195 200 205

Glu Asp Glu Gly Val Tyr Gln Cys His Ala Ala Asn Met Val Gly Glu
210 215 220

Ala Glu Ser His Ser Thr Val Thr Val Leu Asp Leu Ser Lys Tyr Arg
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Ser Phe His Phe Pro Ala Pro Asp Asp Arg Met
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<210> 11

<211> 103

<212> PRT

<213> Homo sapiens

<400> 11

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Val Arg Ala Val Pro Thr Pro Val Ile Thr Trp Arg Lys Val Thr Lys
35 40 45

Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu Leu Pro Gly Asp His Val
50 55 60

Asn Ile Ala Val Gln Val Arg Gly Pro Ser Asp His Glu Ala Thr
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Gln Cys His Ala Ala Asn Met
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HYS-38CIP sequence listing

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<211> 390

<212> PRT

<213> Homo sapiens

<400> 12

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Gly Arg Arg Pro Lys Cys Gly Pro Cys Arg Pro Glu Gly Cys Pro Ala
35 40 45

Pro Ala Pro Cys Pro Ala Pro Gly Ile Ser Ala Leu Asp Glu Cys Gly
50 55 60

Cys Cys Ala Arg Cys Leu Gly Ala Glu Gly Ala Ser Cys Gly Gly Arg
65 70 75 80

Ala Gly Gly Arg Cys Gly Pro Gly Leu Val Cys Ala Ser Gln Ala Ala
85 90 95

Gly Ala Ala Pro Glu Gly Thr Gly Leu Cys Val Cys Ala Gln Arg Gly
100 105 110

Thr Val Cys Gly Ser Asp Gly Arg Ser Tyr Pro Ser Val Cys Ala Leu
115 120 125

Arg Leu Arg Ala Arg His Thr Pro Arg Ala His Pro Gly His Leu His
130 135 140

Lys Ala Arg Asp Gly Pro Cys Glu Phe Val Pro Ile Thr Arg Phe Tyr
145 150 155 160

Asn Cys Phe Pro Gln Pro Leu Ile His Arg Gln Phe Ser Leu Ser Pro
165 170 175

Asp Arg Arg Gln Ser Glu Thr Leu Ser Lys Lys Lys Lys Lys Glu
180 185 190

Glu Glu Glu Glu Glu Glu Gly Glu Glu Glu Lys Glu Glu Glu
Page 10

HYS-38CIP sequence listing
195 200 205

Gly Cys Lys Ser Asn Phe Gln His Thr Ile Asn Phe Lys Glu Ile Ser
210 215 220

Glu Gly Phe Gly Lys Ile Phe Ser Phe Gln Pro Ser Met Ile Asp Ile
225 230 235 240

Ile Asp Glu Ala Ser Thr Leu His Val Ala Gln His Ala Val Val Leu
245 250 255

Asp Ala Arg Val Ala Glu Leu Leu Ser Asn Ala Ala Pro Val Val Val
260 265 270

Val Pro Pro Arg Ser Val His Asn Val Thr Gly Ala Gln Val Gly Leu
275 280 285

Ser Cys Glu Val Arg Ala Val Pro Thr Pro Val Ile Thr Trp Arg Lys
290 295 300

Val Thr Lys Ser Pro Glu Gly Thr Gln Ala Leu Glu Glu Leu Pro Gly
305 310 315 320

Asp His Val Asn Ile Ala Val Gln Val Arg Gly Gly Pro Ser Asp His
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Glu Ala Thr Ala Trp Ile Leu Val Ser Asp Leu His His Cys Leu Lys
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Ala Leu Pro Thr Tyr Ser Tyr Ser Ser Thr Leu Ser Pro Ser Gln Val
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<210> 13

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<212> PRT

<213> Mus musculus

<400> 13

HYS-38CIP sequence listing

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20 25 30

Cys Gln Gln Asp Arg Cys Pro Ala Pro Ser Pro Cys Pro Ala Pro Trp
35 40 45

Ile Ser Ala Arg Asp Glu Cys Gly Cys Cys Ala Arg Cys Leu Gly Ala
50 55 60

Glu Gly Ala Ser Cys Gly Gly Pro Val Gly Ser Arg Cys Gly Pro Gly
65 70 75 80

Leu Val Cys Ala Ser Arg Ala Ser Gly Thr Ala Pro Glu Gly Thr Gly
85 90 95

Leu Cys Val Cys Ala Gln Arg Gly Ala Val Cys Gly Ser Asp Gly Arg
100 105 110

Ser Tyr Ser Ser Ile Cys Ala Leu Arg Leu Arg Ala Arg His Ala Pro
115 120 125

Arg Ala His His Gly His Leu His Lys Ala Arg Asp Gly Pro Cys Glu
130 135 140

Phe Ala Pro Val Val Leu Met Pro Pro Arg Asp Ile His Asn Val Thr
145 150 155 160

Gly Thr Gln Val Phe Leu Ser Cys Glu Val Lys Ala Val Pro Thr Pro
165 170 175

Val Ile Thr Trp Lys Lys Val Lys His Ser Pro Glu Gly Thr Glu Gly
180 185 190

Leu Glu Glu Leu Pro Gly Asp His Val Asn Ile Ala Val Gln Val Arg
195 200 205

Gly Gly Pro Ser Asp His Glu Thr Thr Ser Trp Ile Leu Ile Asn Pro
210 215 220

Leu Arg Lys Glu Asp Glu Gly Val Tyr His Cys His Ala Ala Asn Ala
225 230 235 240

Ile Gly Glu Ala Gln Ser His Gly Thr Val Thr Val Leu Asp Leu Asn
245 250 255

HYS-38CIP sequence listing

Arg Tyr Lys Ser Leu Tyr Ser Ser Val Pro Gly Asp
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<210> 14

<211> 264

<212> PRT

<213> Homo sapiens

<400> 14

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20 25 30

Glu Pro Ala Ser Cys Pro Pro Leu Pro Pro Leu Gly Cys Leu Leu Gly
35 40 45

Glu Thr Arg Asp Ala Cys Gly Cys Cys Pro Met Cys Ala Arg Gly Glu
50 55 60

Gly Glu Pro Cys Gly Gly Gly Ala Gly Arg Gly Tyr Cys Ala Pro
65 70 75 80

Gly Met Glu Cys Val Lys Ser Arg Lys Arg Arg Lys Gly Lys Ala Gly
85 90 95

Ala Ala Ala Gly Gly Pro Gly Val Ser Gly Val Cys Val Cys Lys Ser
100 105 110

Arg Tyr Pro Val Cys Gly Ser Asp Gly Thr Thr Tyr Pro Ser Gly Cys
115 120 125

Gln Leu Arg Ala Ala Ser Gln Arg Ala Glu Ser Arg Gly Glu Lys Ala
130 135 140

Ile Thr Gln Val Ser Lys Gly Thr Cys Glu Gln Gly Pro Ser Ile Val
145 150 155 160

Thr Pro Pro Lys Asp Ile Trp Asn Val Thr Gly Ala Gln Val Tyr Leu
165 170 175

Ser Cys Glu Val Ile Gly Ile Pro Thr Pro Val Leu Ile Trp Asn Lys
180 185 190

HYS-38CIP sequence listing

Val Lys Arg Gly His Tyr Gly Val Gln Arg Thr Glu Leu Leu Pro Gly
195 200 205

Asp Arg Asp Asn Leu Ala Ile Gln Thr Arg Gly Gly Pro Glu Lys His
210 215 220

Glu Val Thr Gly Trp Val Leu Val Ser Pro Leu Ser Lys Glu Asp Ala
225 230 235 240

Gly Glu Tyr Glu Cys His Ala Ser Asn Phe Gln Gly Gln Ala Ser Ala
245 250 255

Ser Ala Lys Ile Thr Val Val Asp
260